

Figure 2: Mouse Genome; One Mistmatch; Truncation to Length "L"

Theory and Measurements

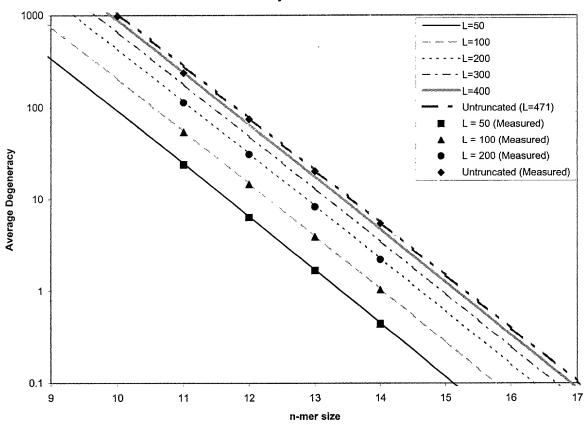
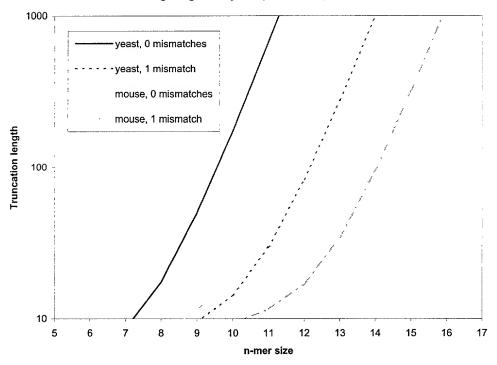
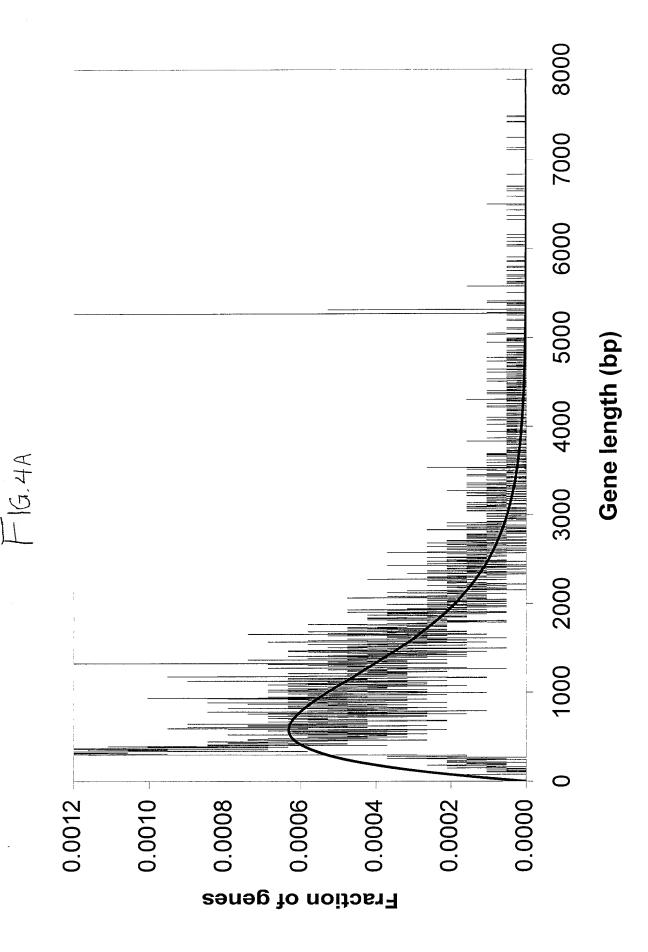
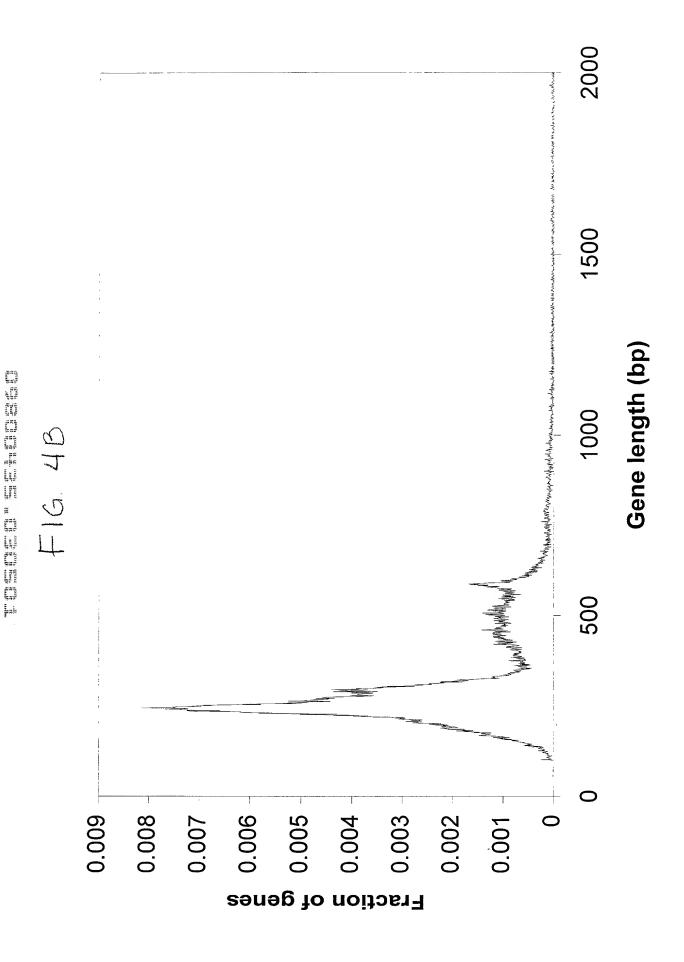


Figure 3: Relationships between parameter values such that average degeneracy = 1 (theoretical predictions)

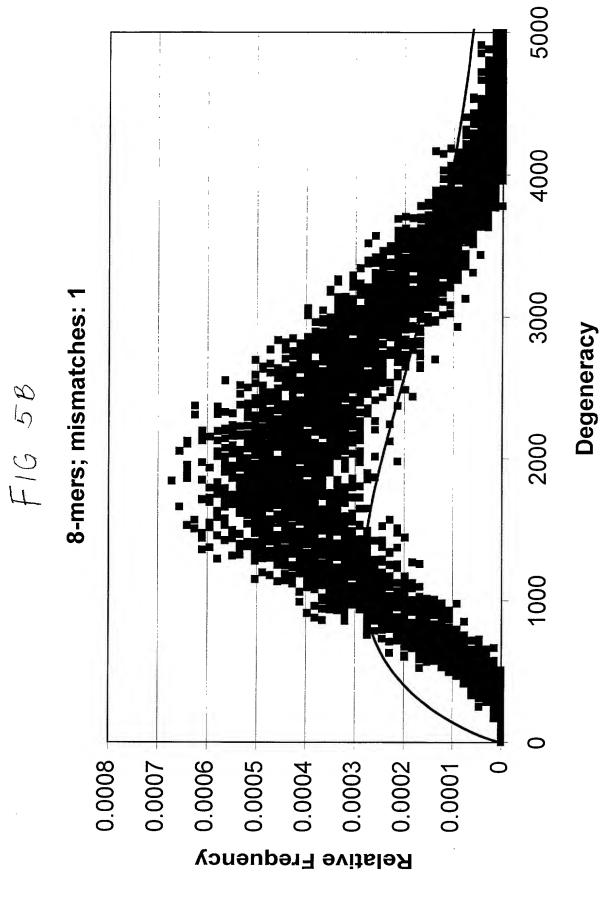


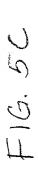


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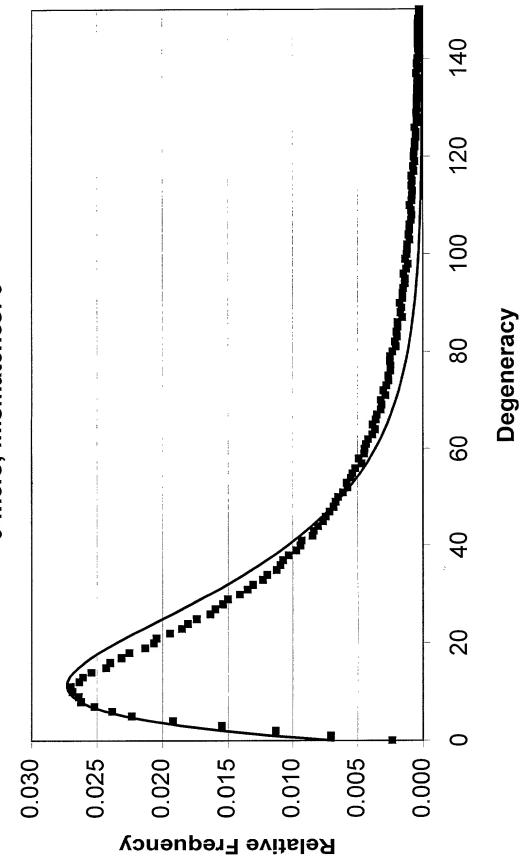


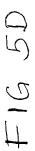
Degeneracy

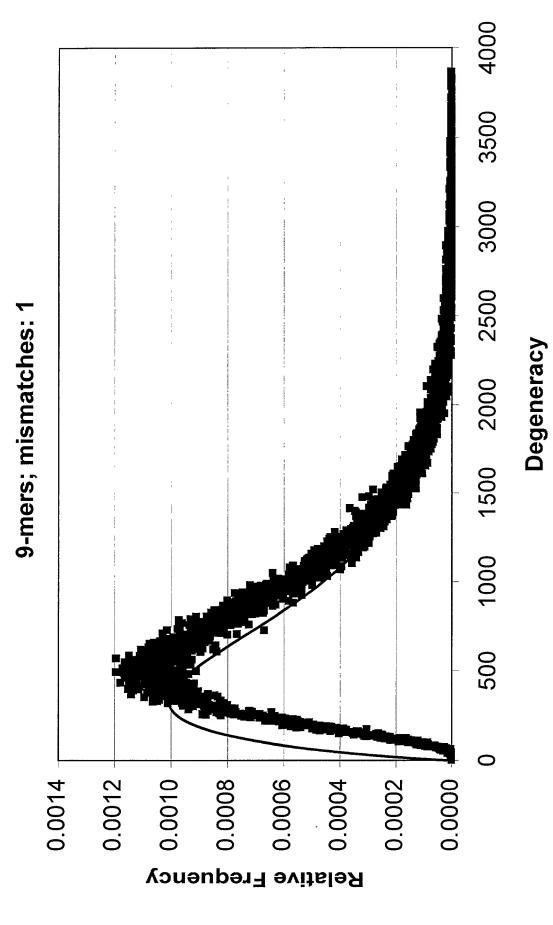


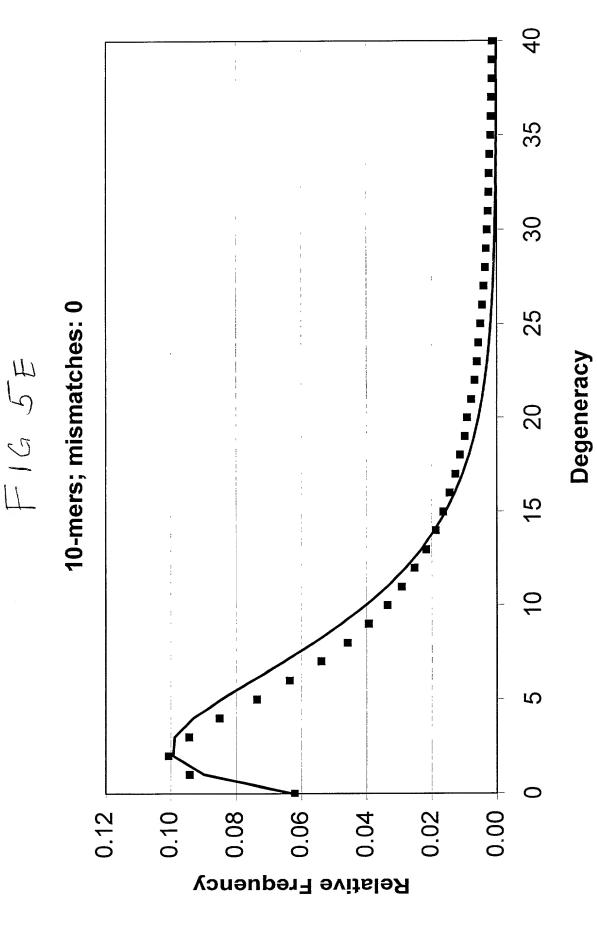


9-mers; mismatches: 0



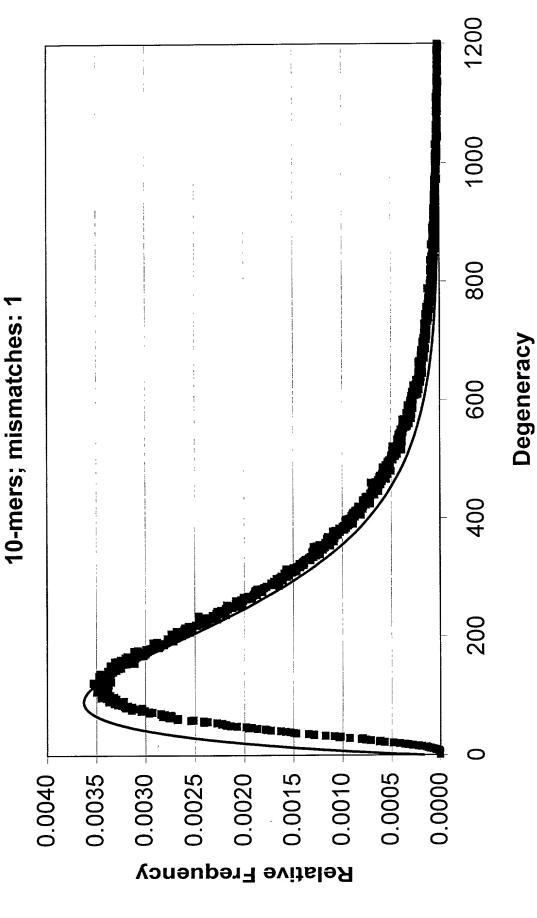


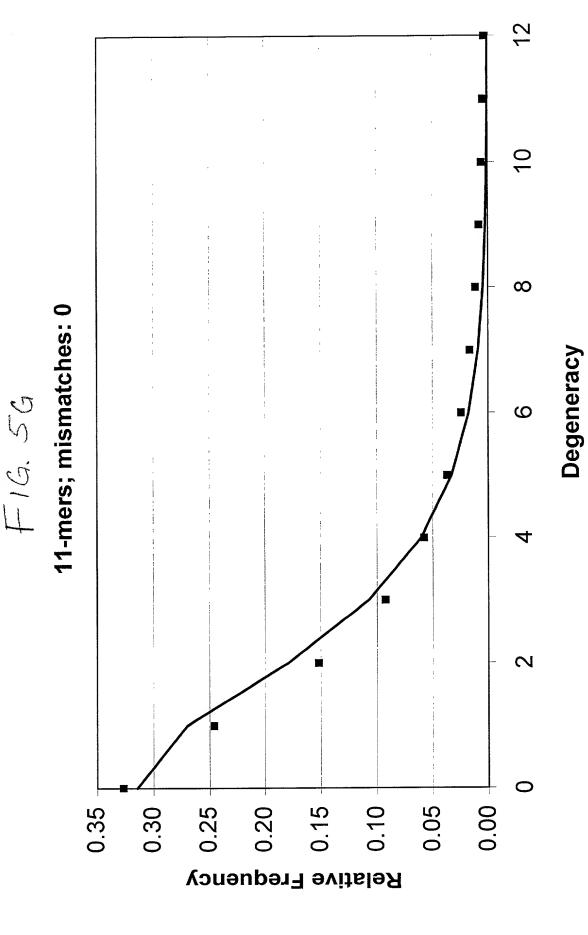


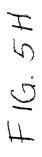


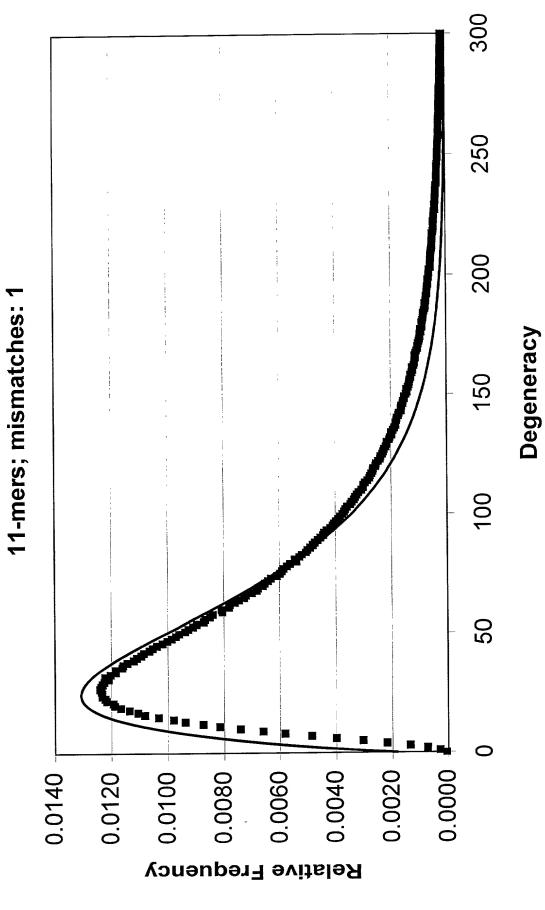


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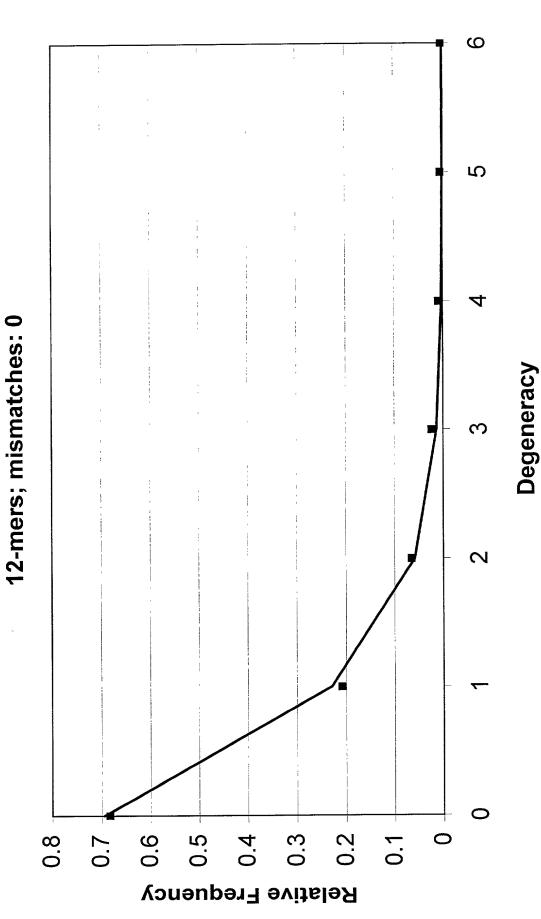




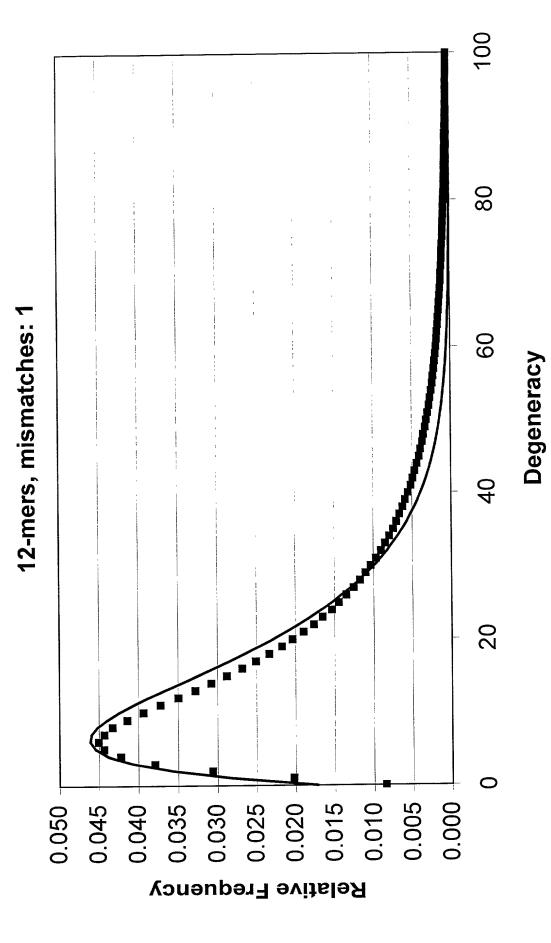


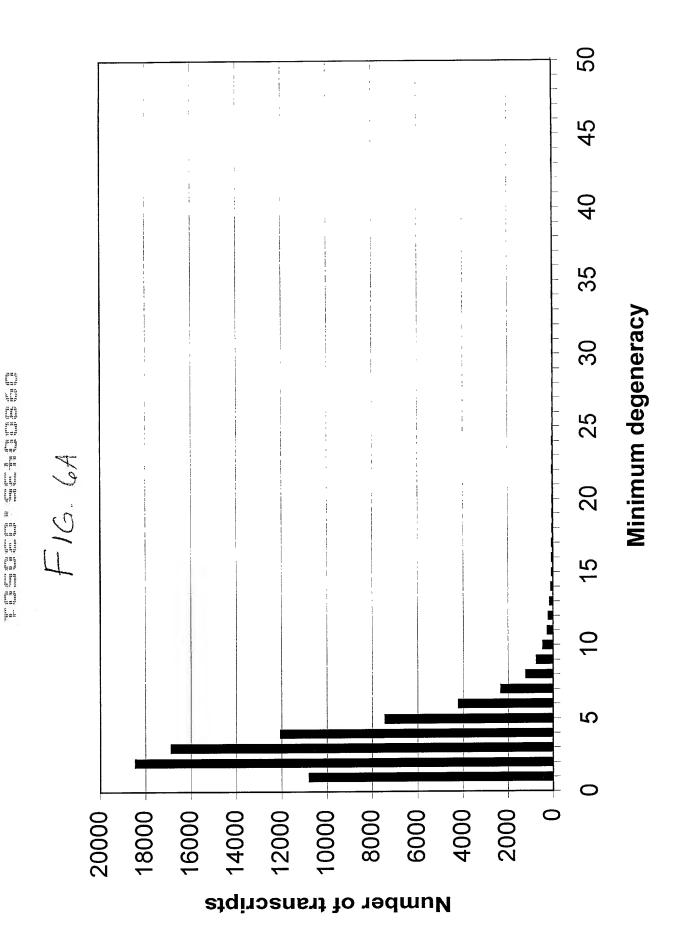


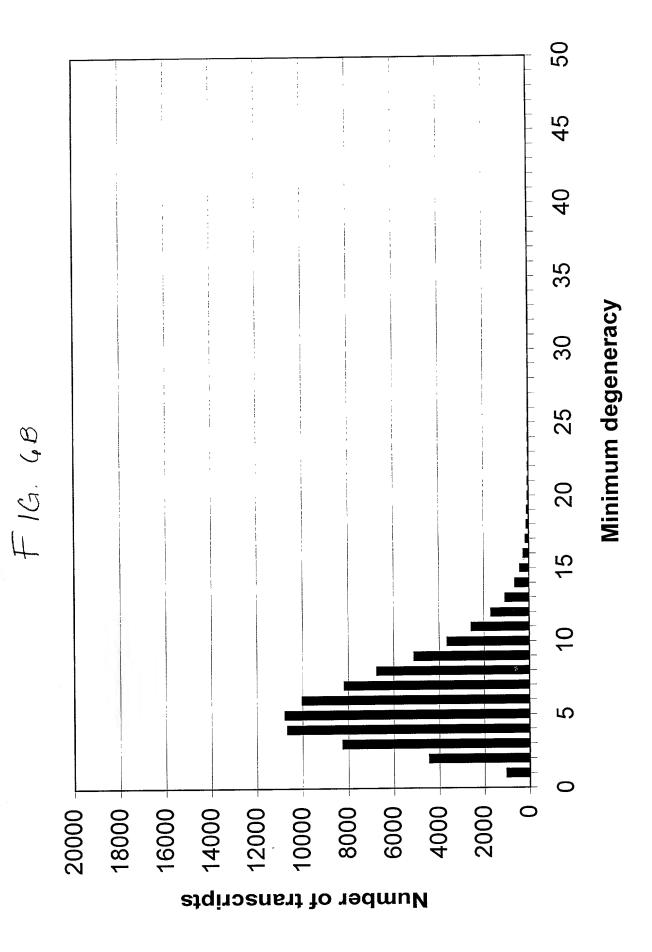




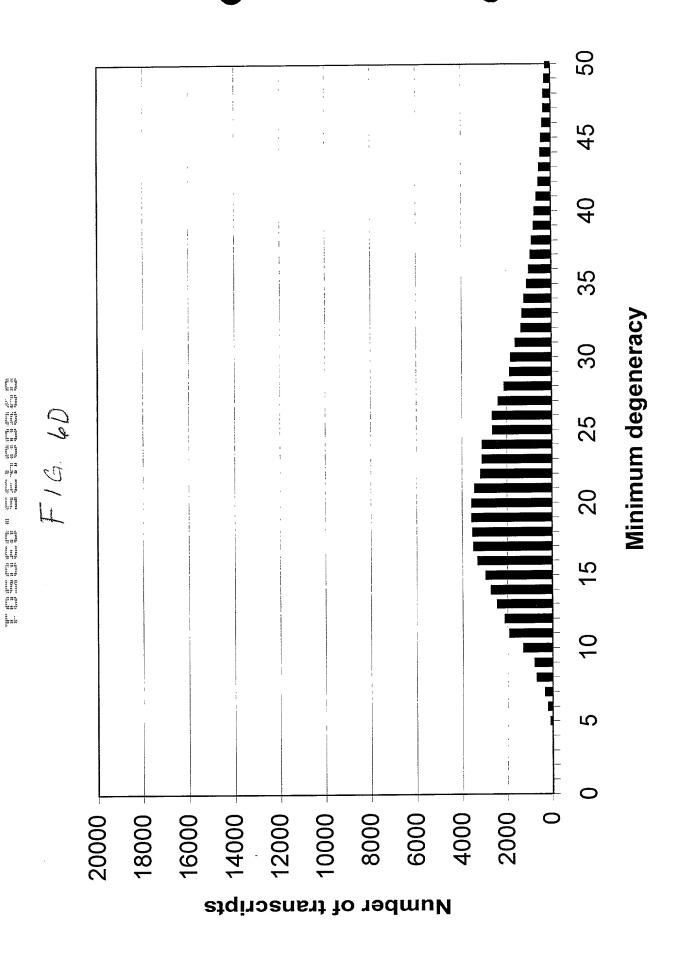


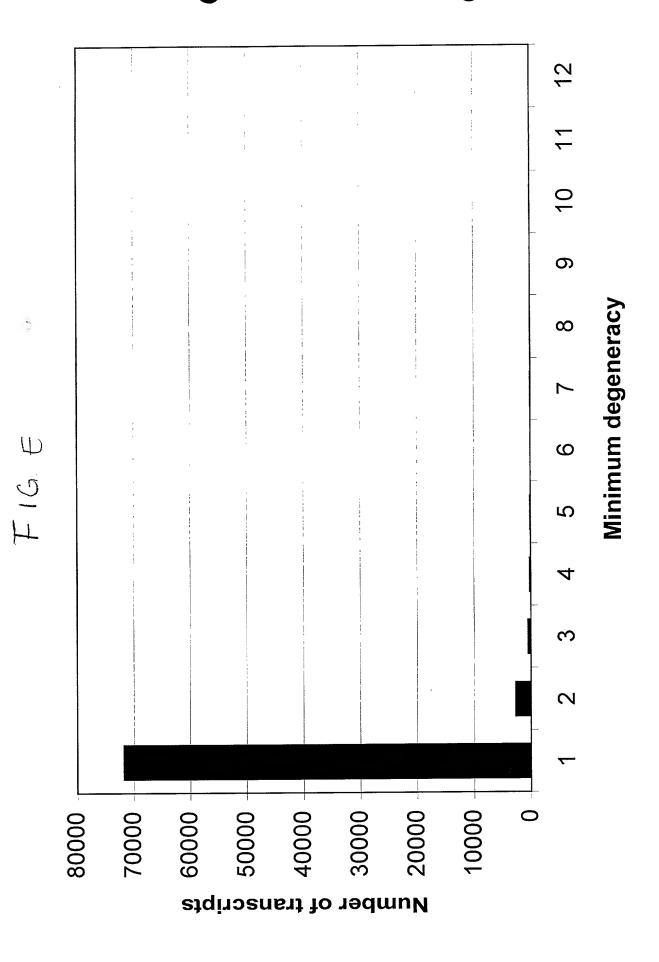


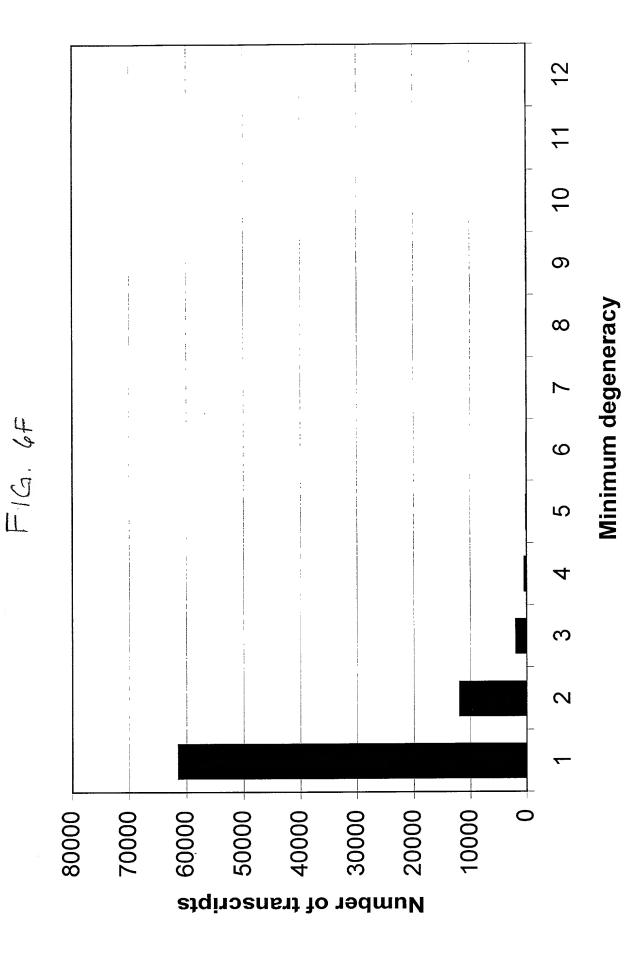




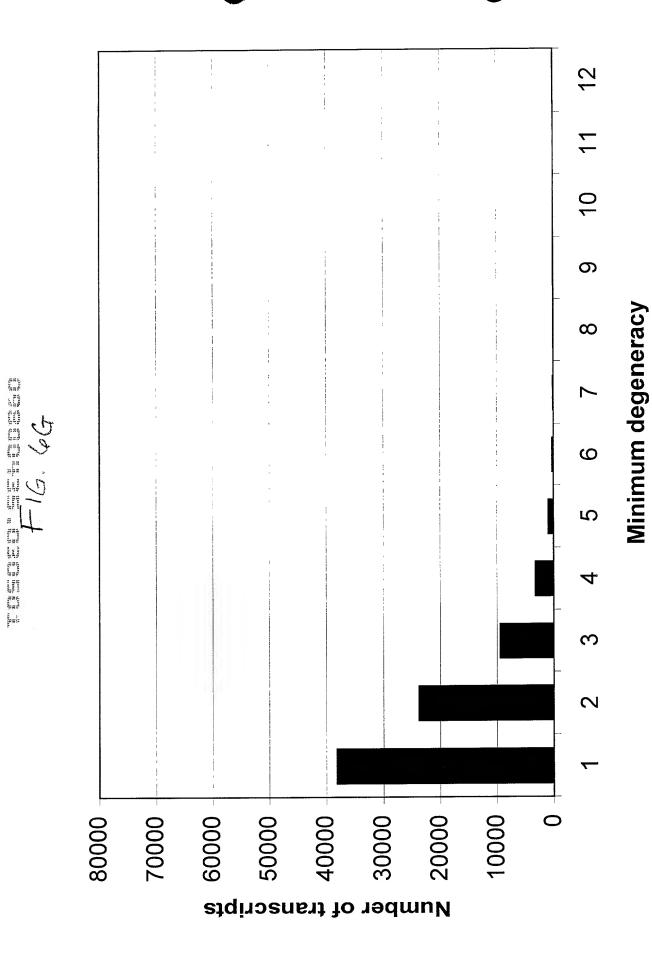
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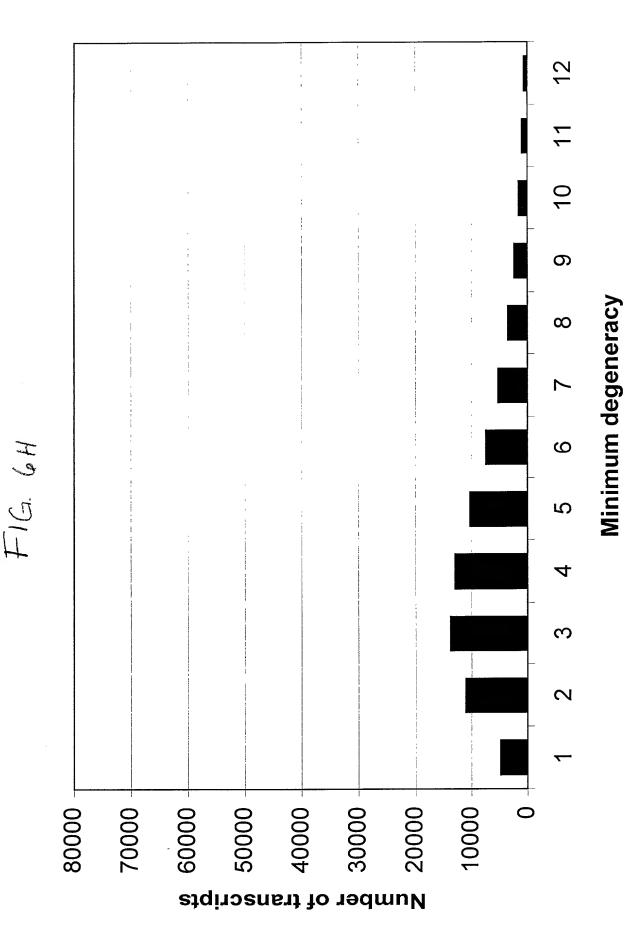


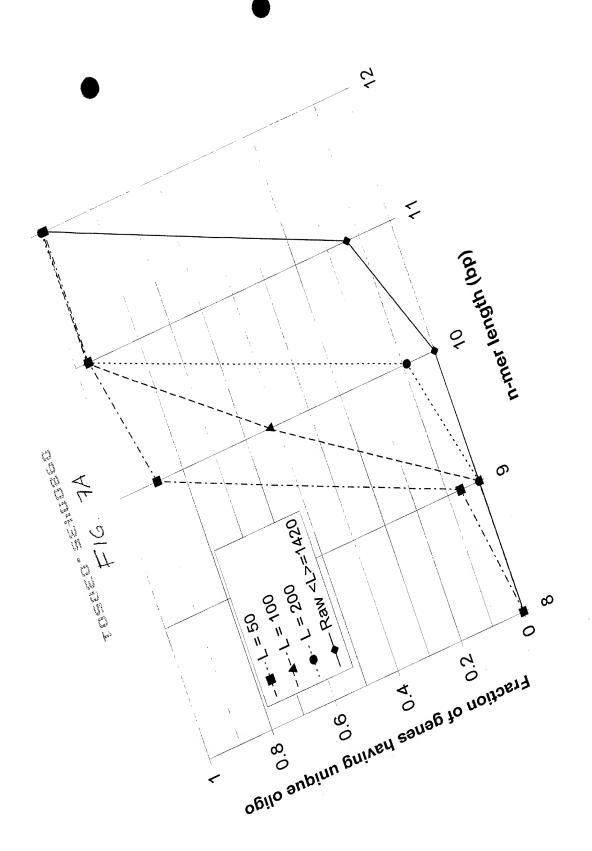




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n-mer length (bp)

the first from 1979, from 1979, the first from 1979